



#6

## SEQUENCE LISTING

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<120> HUMANIZED ANTIBODIES THAT RECOGNIZE VEROTOXIN II AND  
CELL LINE PRODUCING SAME

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<140> 09/700851

<141> 2000-11-17

<150> WO 99/59629

<151> 1999-05-19

<150> US 60/086,570

<151> 1998-05-20

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<170> PatentIn Ver. 2.1

<210> 1

<211> 414

<212> DNA

<213> Mus musculus

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<221> CDS

<222> (1)..(414)

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<223> Figure 1(A): Heavy chain variable region of mouse  
antibody VTm1.1 (MuVTm1.1).

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atg aac ttt gtg ctc agc tcg att ttc ctt gcc ctc att tta aaa gga	48
Met Asn Phe Val Leu Ser Ser Ile Phe Leu Ala Leu Ile Leu Lys Gly	
1 5 10 15	

gtc cag tgt gaa gtg cag ctg gtg gag tcg ggg gga ggc tta gtg aag	96
Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Lys	
20 25 30	

cct gga ggg ccc ctg aaa ctc tcc tgt gca gcc tct gga ttc act ttc	144
Pro Gly Gly Pro Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe	
35 40 45	

agt agt tat ggc atg tct tgg gtt cgc cag act ccg gag aag agg ctg	192
Ser Ser Tyr Gly Met Ser Trp Val Arg Gln Thr Pro Glu Lys Arg Leu	
50 55 60	

gag tgg gtc gca acc att agt act ggt ggt agt tac acc tac tac cca	240
Glu Trp Val Ala Thr Ile Ser Thr Gly Gly Ser Tyr Thr Tyr Tyr Pro	
65 70 75 80	

gac agt gtg aag ggt cga ttc acc atc tcc aga gac aat gcc aag aac 288  
 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn  
                     85                                    90                                    95

gcc ctg tat ctg caa atg agc agt ctg agg tct gag gac acg gcc ata 336  
 Ala Leu Tyr Leu Gln Met Ser Ser Leu Arg Ser Glu Asp Thr Ala Ile  
                     100                                    105                                    110

tat tac tgt gca aga cgg ggg gac gca tgg ggt aac ttg gac tac tgg 384  
 Tyr Tyr Cys Ala Arg Arg Gly Asp Ala Trp Gly Asn Leu Asp Tyr Trp  
                     115                                    120                                    125

ggt caa gga acc tct gtc acc gtc tcc tca 414  
 Gly Gln Gly Thr Ser Val Thr Val Ser Ser  
                     130                                    135

<210> 2

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<212> PRT

<213> Mus musculus

<220>

<223> Figure 1(A): Heavy chain variable region of mouse  
 antibody VTm1.1 (MuVTm1.1).

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Met Asn Phe Val Leu Ser Ser Ile Phe Leu Ala Leu Ile Leu Lys Gly  
                     1                                    5                                    10                                    15

Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Lys  
                     20                                    25                                    30

Pro Gly Gly Pro Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe  
                     35                                    40                                    45

Ser Ser Tyr Gly Met Ser Trp Val Arg Gln Thr Pro Glu Lys Arg Leu  
                     50                                    55                                    60

Glu Trp Val Ala Thr Ile Ser Thr Gly Gly Ser Tyr Thr Tyr Tyr Pro  
                     65                                    70                                    75                                    80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn  
                     85                                    90                                    95

Ala Leu Tyr Leu Gln Met Ser Ser Leu Arg Ser Glu Asp Thr Ala Ile  
                     100                                    105                                    110

Tyr Tyr Cys Ala Arg Arg Gly Asp Ala Trp Gly Asn Leu Asp Tyr Trp  
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Gly Gln Gly Thr Ser Val Thr Val Ser Ser  
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<212> DNA

<213> Mus musculus

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 <221> CDS  
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 <223> Figure 1(B): Light chain variable region of mouse  
 VTm1.1 antibody (MuVTm1.1).

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<400> 3
atg gtt ttc aca cct cag ata ctt gga ctt atg ctt ttt tgg att tca 48
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      1              5              10              15

gcc tcc aga ggt gat gtt gtg cta act cag tct cca gcc acc ctg tct 96
Ala Ser Arg Gly Asp Val Val Leu Thr Gln Ser Pro Ala Thr Leu Ser
      20              25              30

gtg act cca gga gat agc gtc agt ctt tcc tgc agg gcc agt caa act 144
Val Thr Pro Gly Asp Ser Val Ser Leu Ser Cys Arg Ala Ser Gln Thr
      35              40              45

att agc aac aac cta cac tgg tat caa cac aaa tca cat gag tct cca 192
Ile Ser Asn Asn Leu His Trp Tyr Gln His Lys Ser His Glu Ser Pro
      50              55              60

agg ctt ctc atc aag tct gct tcc cag tcc atc tct ggg atc ccc tcc 240
Arg Leu Leu Ile Lys Ser Ala Ser Gln Ser Ile Ser Gly Ile Pro Ser
      65              70              75              80

agg ttc agt ggc agt gga tca ggg aca gat ttc act ctc agt atc aac 288
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn
      85              90              95

agt gtg gaa act gaa gat ttt gga atg tat ttc tgt caa cag agt tac 336
Ser Val Glu Thr Glu Asp Phe Gly Met Tyr Phe Cys Gln Gln Ser Tyr
      100              105              110

agc tgg ccg ctc acg ttc ggt gct ggg acc aag ctg gag ctg aaa 381
Ser Trp Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys
      115              120              125

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<210> 4  
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 <213> Mus musculus

<220>  
 <223> Figure 1(B): Light chain variable region of mouse  
 VTm1.1 antibody (MuVTm1.1).

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<400> 4
Met Val Phe Thr Pro Gln Ile Leu Gly Leu Met Leu Phe Trp Ile Ser
      1              5              10              15

Ala Ser Arg Gly Asp Val Val Leu Thr Gln Ser Pro Ala Thr Leu Ser
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Val Thr Pro Gly Asp Ser Val Ser Leu Ser Cys Arg Ala Ser Gln Thr
      35              40              45

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Tyr Tyr Cys Ala Arg Arg Gly Asp Ala Trp Gly Asn Leu Asp Tyr Trp  
 115 120 125

ggt caa gga acc tta gtc acc gtc tcc tca  
 Gly Gln Gly Thr Leu Val Thr Val Ser Ser  
 130 135

414

<210> 6  
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 <212> PRT  
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 <223> Figure 2(A): Heavy chain variable region of  
 humanized VTm1.1 antibody (HuVTm1.1).

<400> 6  
 Met Asn Phe Val Leu Ser Ser Ile Phe Leu Ala Leu Ile Leu Lys Gly  
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 Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln  
 20 25 30  
 Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe  
 35 40 45  
 Ser Ser Tyr Gly Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu  
 50 55 60  
 Glu Trp Val Ala Thr Ile Ser Thr Gly Gly Ser Tyr Thr Tyr Tyr Pro  
 65 70 75 80  
 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn  
 85 90 95  
 Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val  
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 Tyr Tyr Cys Ala Arg Arg Gly Asp Ala Trp Gly Asn Leu Asp Tyr Trp  
 115 120 125  
 Gly Gln Gly Thr Leu Val Thr Val Ser Ser  
 130 135

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 <212> DNA  
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<220>  
 <221> CDS  
 <222> (1)..(381)  
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 <223> igure 2(B): Light chain variable region of  
 humanized VTm1.1 antibody (HuVTm1.1) .

<400> 7

atg gtt ttc aca cct cag ata ctt gga ctt atg ctt ttt tgg att tca	48
Met Val Phe Thr Pro Gln Ile Leu Gly Leu Met Leu Phe Trp Ile Ser	
1 5 10 15	
gcc tcc aga ggt gaa att gtg cta act cag tct cca gcc acc ctg tct	96
Ala Ser Arg Gly Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser	
20 25 30	
gtg tct cca gga gaa aga gcc act ctt tcc tgc agg gcc agt caa act	144
Val Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Thr	
35 40 45	
att agc aac aac cta cac tgg tat caa caa aaa cca ggt cag gct cca	192
Ile Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro	
50 55 60	
agg ctt ctc atc aag tct gct tcc cag tcc atc tct ggg ata ccc gcc	240
Arg Leu Leu Ile Lys Ser Ala Ser Gln Ser Ile Ser Gly Ile Pro Ala	
65 70 75 80	
agg ttc agt ggc agt gga tca ggg aca gat ttc act ctc act atc agc	288
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser	
85 90 95	
agt ctg gaa tct gaa gat ttt gca gtg tat tac tgt caa cag agt tac	336
Ser Leu Glu Ser Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Ser Tyr	
100 105 110	
agt tgg ccg ctc acg ttc ggt caa ggg acc aag gtg gag atc aaa	381
Ser Trp Pro Leu Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys	
115 120 125	

<210> 8

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<223> igure 2(B): Light chain variable region of  
humanized VTm1.1 antibody (HuVTm1.1) .

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Met Val Phe Thr Pro Gln Ile Leu Gly Leu Met Leu Phe Trp Ile Ser	
1 5 10 15	
Ala Ser Arg Gly Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser	
20 25 30	
Val Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Thr	
35 40 45	
Ile Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro	
50 55 60	
Arg Leu Leu Ile Lys Ser Ala Ser Gln Ser Ile Ser Gly Ile Pro Ala	
65 70 75 80	
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser	
85 90 95	

Ser	Leu	Glu	Ser	Glu	Asp	Phe	Ala	Val	Tyr	Tyr	Cys	Gln	Gln	Ser	Tyr
			100					105					110		
Ser	Trp	Pro	Leu	Thr	Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile	Lys	
		115					120					125			